

OIPE

RAW SEQUENCE LISTING

DATE: 10/29/2001

PATENT APPLICATION: US/09/974,712

TIME: 15:40:30

Input Set : A:\LEX-0251-USA SEQLIST.txt

Output Set: N:\CRF3\10292001\I974712.raw

4 <110> APPLICANT: Friddle, Carl Johan

5 Hilbun, Erin

6 Gerhardt, Brenda

7 Turner, C. Alexander Jr.

9 <120> TITLE OF INVENTION: Novel Human Ion Channel Protein and Polynucleotides Encoding
the Same

11 <130> FILE REFERENCE: LEX-0251-USA

C--> 13 <140> CURRENT APPLICATION NUMBER: US/09/974,712

C--> 13 <141> CURRENT FILING DATE: 2001-10-10

13 <150> PRIOR APPLICATION NUMBER: US 60/239,623

14 <151> PRIOR FILING DATE: 2000-10-10

16 <160> NUMBER OF SEQ ID NOS: 3

18 <170> SOFTWARE: FastSEQ for Windows Version 4.0

20 <210> SEQ ID NO: 1

21 <211> LENGTH: 1371

22 <212> TYPE: DNA

23 <213> ORGANISM: homo sapiens

25 <400> SEQUENCE: 1

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28	gaccacagcg	gccgcggccg	cttctacgac	gacgcgcgc	gcgagtattt	cttcgaccgg	180
29	caccggccca	gcttcgacgc	cgtgctctac	tactaccagt	ccggtgggcg	gctgcggcgg	240
30	ccggcgacg	tgccgctcga	cgtcttcctg	gaagaggtag	ccttctacgg	gctgggcgcg	300
31	gcggccctgg	cacgcctgcg	cgaggacgag	ggctgcccgg	tgccgcccga	gcgccccctg	360
32	ccccgcgcg	ccttcgcccg	ccagctgtgg	ctgcttttcg	agtttcccga	gagctctcag	420
33	gcgcgcgcg	tgctgcgcgt	agtctccgtg	ctggtcaccc	tcgtctccat	cgtcgtcttc	480
34	tgccctcaga	cgctgcctga	cttcgcgcac	gaccgcgacg	gcacggggct	tgctgctgca	540
35	gccgcagccg	gcccgttccc	cgctcggctg	aatggctcca	gccaaatgcc	tggaaatcca	600
36	ccccgcctgc	ccttcaatga	cccgttcttc	gtggtggaga	cgctgtgtat	ttgttggttc	660
37	tcctttgagc	tgctggtagc	cctcctggtc	tgtccaagca	aggctatott	cttcaagaac	720
38	gtgatgaacc	tcacgcattt	tgtggctatc	cttcctact	ttgtggcact	gggcaccgag	780
39	ctggcccggc	agcgaggggt	gggccagcag	gccatgtcac	tggccatcct	gagagtcac	840
40	cgattggtag	gtgtcttcgc	catcttcaag	ctgtcccggc	actcaaagg	cctgcaaata	900
41	ttgggccaga	cgttcggggc	ctccatgcgt	gagctggggc	tcctcatctt	tttctcttc	960
42	atcggtgtgg	tcctcttttc	cagcgccgtc	tactttgccg	aagttgaccg	ggtggactcc	1020
43	catttcaacta	gcacccctga	gtccttctg	tgggcgtag	tcacatgac	tacagttggc	1080
44	tatggagaca	tggcacccgt	cactgtgggt	ggcaagatag	tgggctctct	gtgtgccatt	1140
45	gcgggcgtgc	tgactatttc	cctgccagtg	cccgtcattg	tctccaattt	cagctacttt	1200
46	tatcaccggg	agacagaggg	cgaagaggct	gggatgttca	gccatgtgga	catgcagcct	1260
47	tgtggccccc	tggagggcaa	ggccaatggg	gggctggtgg	acggggaggt	acctgagcta	1320
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50 <210> SEQ ID NO: 2

51 <211> LENGTH: 456

52 <212> TYPE: PRT

53 <213> ORGANISM: homo sapiens

55 <400> SEQUENCE: 2

56 Met Glu Pro Arg Cys Pro Pro Pro Cys Gly Cys Cys Glu Arg Leu Val

57 1

5

10

15

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58 Leu Asn Val Ala Gly Leu Arg Phe Glu Thr Arg Ala Arg Thr Leu Gly
59          20          25          30
60 Arg Phe Pro Asp Thr Leu Leu Gly Asp Pro Ala Arg Arg Gly Arg Phe
61          35          40          45
62 Tyr Asp Asp Ala Arg Arg Glu Tyr Phe Phe Asp Arg His Arg Pro Ser
63          50          55          60
64 Phe Asp Ala Val Leu Tyr Tyr Tyr Gln Ser Gly Gly Arg Leu Arg Arg
65 65          70          75          80
66 Pro Ala His Val Pro Leu Asp Val Phe Leu Glu Glu Val Ala Phe Tyr
67          85          90          95
68 Gly Leu Gly Ala Ala Ala Leu Ala Arg Leu Arg Glu Asp Glu Gly Cys
69          100          105          110
70 Pro Val Pro Pro Glu Arg Pro Leu Pro Arg Arg Ala Phe Ala Arg Gln
71          115          120          125
72 Leu Trp Leu Leu Phe Glu Phe Pro Glu Ser Ser Gln Ala Ala Arg Val
73          130          135          140
74 Leu Ala Val Val Ser Val Leu Val Ile Leu Val Ser Ile Val Val Phe
75 145          150          155          160
76 Cys Leu Glu Thr Leu Pro Asp Phe Arg Asp Asp Arg Asp Gly Thr Gly
77          165          170          175
78 Leu Ala Ala Ala Ala Ala Ala Gly Pro Phe Pro Ala Arg Leu Asn Gly
79          180          185          190
80 Ser Ser Gln Met Pro Gly Asn Pro Pro Arg Leu Pro Phe Asn Asp Pro
81          195          200          205
82 Phe Phe Val Val Glu Thr Leu Cys Ile Cys Trp Phe Ser Phe Glu Leu
83          210          215          220
84 Leu Val Arg Leu Leu Val Cys Pro Ser Lys Ala Ile Phe Phe Lys Asn
85 225          230          235          240
86 Val Met Asn Leu Ile Asp Phe Val Ala Ile Leu Pro Tyr Phe Val Ala
87          245          250          255
88 Leu Gly Thr Glu Leu Ala Arg Gln Arg Gly Val Gly Gln Gln Ala Met
89          260          265          270
90 Ser Leu Ala Ile Leu Arg Val Ile Arg Leu Val Arg Val Phe Arg Ile
91          275          280          285
92 Phe Lys Leu Ser Arg His Ser Lys Gly Leu Gln Ile Leu Gly Gln Thr
93          290          295          300
94 Leu Arg Ala Ser Met Arg Glu Leu Gly Leu Leu Ile Phe Phe Leu Phe
95 305          310          315          320
96 Ile Gly Val Val Leu Phe Ser Ser Ala Val Tyr Phe Ala Glu Val Asp
97          325          330          335
98 Arg Val Asp Ser His Phe Thr Ser Ile Pro Glu Ser Phe Trp Trp Ala
99          340          345          350
100 Val Val Thr Met Thr Thr Val Gly Tyr Gly Asp Met Ala Pro Val Thr
101          355          360          365
102 Val Gly Gly Lys Ile Val Gly Ser Leu Cys Ala Ile Ala Gly Val Leu
103          370          375          380
104 Thr Ile Ser Leu Pro Val Pro Val Ile Val Ser Asn Phe Ser Tyr Phe
105 385          390          395          400
106 Tyr His Arg Glu Thr Glu Gly Glu Glu Ala Gly Met Phe Ser His Val

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107                               405                               410                               415
108 Asp Met Gln Pro Cys Gly Pro Leu Glu Gly Lys Ala Asn Gly Gly Leu
109                               420                               425                               430
110 Val Asp Gly Glu Val Pro Glu Leu Pro Pro Pro Leu Trp Ala Pro Pro
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113                               450                               455
115 <210> SEQ ID NO: 3
116 <211> LENGTH: 1792
117 <212> TYPE: DNA
118 <213> ORGANISM: homo sapiens
120 <400> SEQUENCE: 3
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123 gctgctgcga gcggtggtg ctcaacgtgg ccgggtgcg cttcgagacg cgggcgcgca      180
124 cgctgggccc cttcccgac actctgctag gggaccagc gcgcgcggc cgcttctacg      240
125 acgacgcgcg ccgcagtat ttcttcgacc ggcaccggcc cagcttcgac gccgtgctct      300
126 actactacca gtccggtggg cggctgcggc ggccggcgca cgtgccgctc gacgtcttcc      360
127 tggaagaggt ggccttctac gggctgggcy cggcgccctt ggcacgcctg cgcgaggacg      420
128 agggctgccc ggtgccgccc gagcgccccc tgccccgcg cgccttcgcc cgccagctgt      480
129 ggctgctttt cgagtttccc gagagctctc aggcgcgcg cgtgctcgcc gtagtctccg      540
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133 tcgtgggtga gacgctgtgt atttggttgt tctcctttga gctgctggtg cgcctcctgg      780
134 tctgtccaag caaggtatc ttcttcaaga acgtgatgaa cctcatcgat tttgtggcta      840
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150 ttttggggta gtgttgagat aggtttgggtc rtatcatttt gtgagtttcc ta      1792

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VERIFICATION SUMMARY

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L:13 M:270 C: Current Application Number differs, Replaced Current Application No

L:13 M:271 C: Current Filing Date differs, Replaced Current Filing Date